IN THE CLAIMS

- 1. (currently amended) A phage particle displaying on its surface:
 - (a) a dimeric T-cell receptor (dTCR), wherein the dTCR comprises:
 - (i) a first polypeptide in which a TCR α chain variable domain sequence is fused to the N terminus of a TCR α chain constant domain extracellular sequence, and
 - (ii) a second polypeptide wherein a TCR β chain variable domain sequence is fused to the N terminus of a TCR β chain constant deomain extracellular sequence; or
- (b) a single chain TCR (scTCR) polypeptide, wherein the scTCR polypeptide comprises:
 - (i) a first segment comprises a TCR α chain variable domain sequence fused to the N terminus of a TCR α chain constant domain extracellular sequence,
 - (ii) a second segment comprising a TCR b chain variable domain sequence fused to the N terminus of a TCR β chain constant domain extracellular sequence; and
 - (iii) a linker sequence linking either (1) the C terminus of the first segment of the N terminus of the second segment or (2) the C terminus of the second segment to the N terminus of the first segment,

wherein the scTCR or dTCR comprises an interchain disulfide bond linking residues of constant domain sequences.

2-5. (canceled)

- 6. (previously presented) The phage particle of claim 1 wherein the C-terminus of one member of the dTCR or the C-terminus of the scTCR polypeptide is linked by a peptide bond to a surface exposed residue of the phage particle.
 - 7-54. (canceled)
- 55. (withdrawn) A method for the identification of TCRs with a specific characteristic, said method comprising subjecting a diverse library of TCRs displayed on phage particles as claimed in claim 1 to
 - a selection process which selects for said characteristic, and isolating proteinaceous particles which display a TCR having said characteristic, and optionally to an amplification process to multiply the isolated particles and/or
- a screening process which measures said characteristic, identifying those proteinaceous particles which display a TCR with the desired characteristic and isolating these proteinaceous particles, and optionally to an amplification process to multiply the isolated particles.
- 56. (withdrawn) The method of claim 55 wherein the specific characteristic is increased affinity for a TCR ligand.
 - 57. (withdrawn) A method for detecting a TCR ligand complex, comprising steps of:
 - (i) providing the phage particle of claim 1;
 - (ii) contacting the phage particle with a putative ligand complex; and
 - (iii) detecting binding of the phage particle to the putative ligand complexes.
- 58. (withdrawn) The method of claim 57 wherein the putative TCR ligand complex is a peptide-MHC complex.

59. (withdrawn) A method of identifying an inhibitor of the interaction between the phage particle of claim 1 and a TCR-binding ligand, comprising steps of:

contacting the phage particle with a TCR-binding ligand, in the presence of and in the absence of a test compound, and

determining whether the presence of the test compound reduces binding of the phage particle to the TCR-binding ligand, whereby reduced binding identifies the test compound as an inhibitor of the interaction between the phage particle and the TCR-binding ligand.

60-85. (canceled)

- 86. (previously presented) The phage particle of claim 1 wherein the interchain disulfide bond has no equivalent in native T cell receptors.
- 87. (previously presented) The phage particle of claim 6 wherein the interchain disulfide bond has no equivalent in native T cell receptors.
- 88. (withdrawn) The phage particle of claim 1 wherein the interchain disulfide bond is between cysteine residues substituted for Thr 48 of exon 1 of TRAC*01 and Ser 57 of exon 1 of TRBC1*-1 or TRBC2*01 or the non-human equivalent thereof.
- 89. (withdrawn) The phage particle of claim 1 which is a filamentous phage and which displays on its surface a dTCR polypeptide pair comprising:
 - a first polypeptide wherein a sequence corresponding to a TCR α chain variable domain sequence is fused to the N terminus of a sequence corresponding to a TCR α chain constant domain extracellular sequence; and

a second polypeptide wherein a sequence corresponding to a TCR β chain variable domain sequence is fused to the N terminus of a sequence corresponding to a TCR β chain constant domain extracellular sequence,

wherein the first and second polypeptides are linked by a disulfide bond between cysteine residues substituted for Thr 48 of exon 1 of TRAC*01 and Ser 57 of exon 1 of TRBC1*-1 or TRBC2*01 or the non-human equivalent thereof.